AMENDMENT TO THE SPECIFICATION

Please replace paragraph at page 7, line 1, with the following amended paragraph.

X2 is selected from the group consisting of Glu, Asp, Lys, and Val;

X3 is selected from the group consisting of Gly and Ala;

X4 is Pro;

X5 is selected from the group consisting of Thr and Ser;

X6 is selected from the group consisting of Leu, Ile, Val, Ala, and Phe;

X7 is selected from the group consisting of Arg and Lys;

X8 is selected from the group consisting of Gln, Asn, and Glu;

X9 is selected from the group consisting of Trp, Tyr, Cys, Ala, and Phe;

X10 is selected from the group consisting of Leu, Ile, Val, Ala, Phe, Met, and

Lys;

L1 is a linker as described herein; and

n is 0 or 1;

and physiologically acceptable salts thereof.

Please replace Table I in the specification beginning at page 46, line 16 with the following amended Table I.

TABLE I

	TABLET	D 14'
Cor	mound	Relative Potency
(EC ₅₀	npound	1 Otelley
(EC 50)	TPO	4.0
TMD.		1.0
TMP monomer (SEQ ID NO: 1) TMP C-C dimer (SEQ ID NO: 2)		3.5
	$(Gly)_n$ -TMP:	3.3
1	$\mathbf{n} = 0$	4.5
2	n = 0 $n = 1$	4.0
3	n=2	4.0
4	n = 3	4.0
5	n = 4	4.0
6	n = 5	4.0
7	n = 6	4.0
8	n = 7	4.0
9	n = 8	4.5
10	n = 9	4.0
11	n = 10	4.0
12	n = 10 $n = 14$	4.0
13	TMP-GPNG-TMP (SEQ ID NO. 10)	3.0
14	IEGPTLRQCLAARA-GGGGGGGG-IEGPTLRQCLAARA	
17		0.5
	(SEQ ID NO. 11)	
15	IEGPTLRQCLAARA-GGGGGGGG-IEGPTLRQCLAARA	0.5
1,5	(SEQ ID NO. 12)	0.5
16	IEGPTLRQALAARA-GGGGGGGG-IEGPTLRQALAARA	0.5
10	(SEQ ID NO. 13)	0.5
17a	TMP-GGGKGGGG-TMP (SEQ ID NO. 14)	4.0
17b	TMP-GGGK(BrAc)GGGG-TMP (SEQ ID NO. 15)	ND
18	TMP-GGGCGGGG-TMP (SEQ ID NO. 16)	4.0
19	TMP-GGGK(PEG)GGGG-TMP (SEQ ID NO. 17)	5.0
20	TMP-GGGC(PEG)GGGG-TMP (SEQ ID NO. 18)	5.0
21	TMP-GGGNGSGG-TMP (SEQ ID NO. 19)	4.0
22	TMP-GGGCGGGG-TMP (SEQ ID NO. 20)	4.0
	TMP-GGGCGGGG-TMP	•